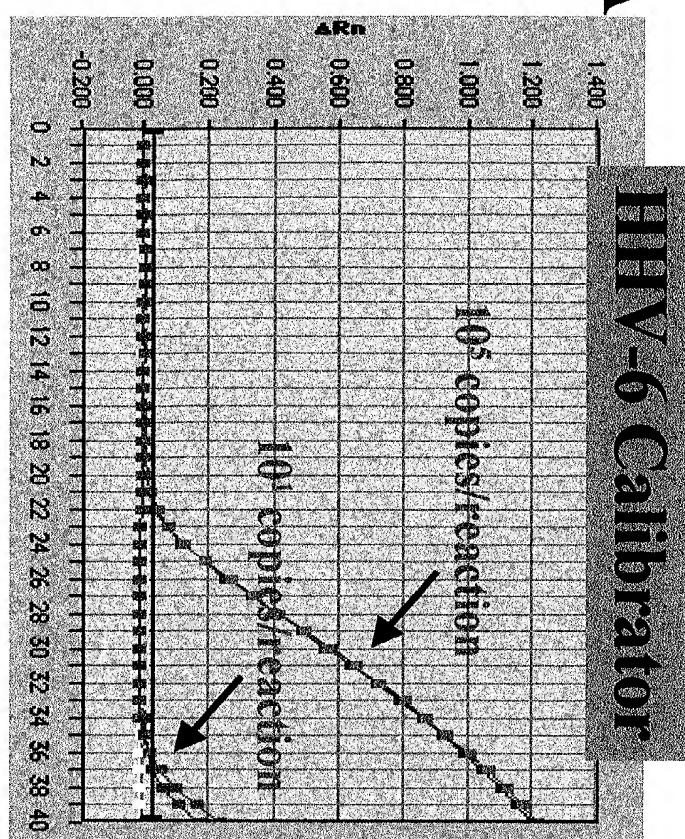
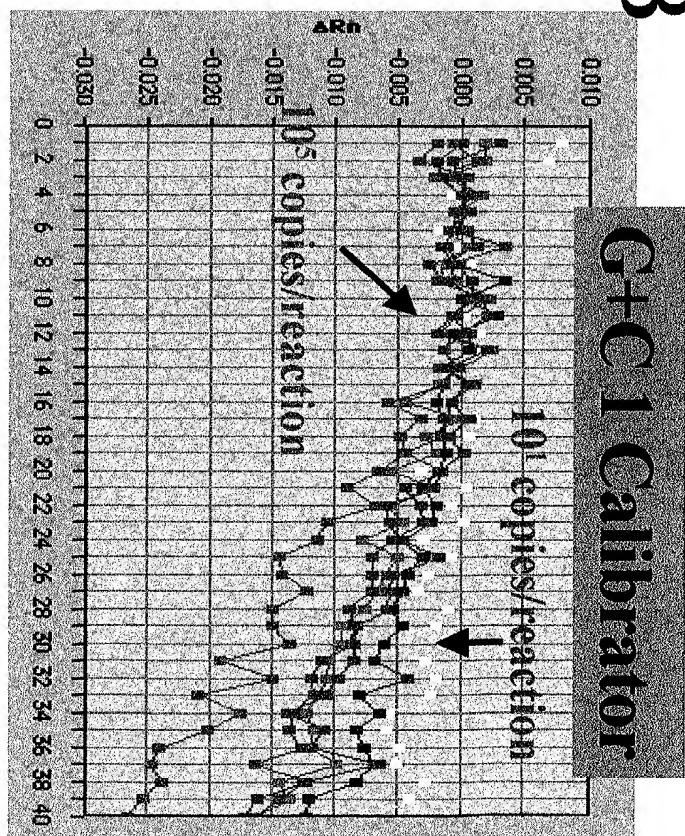


Myers S. Wolitz

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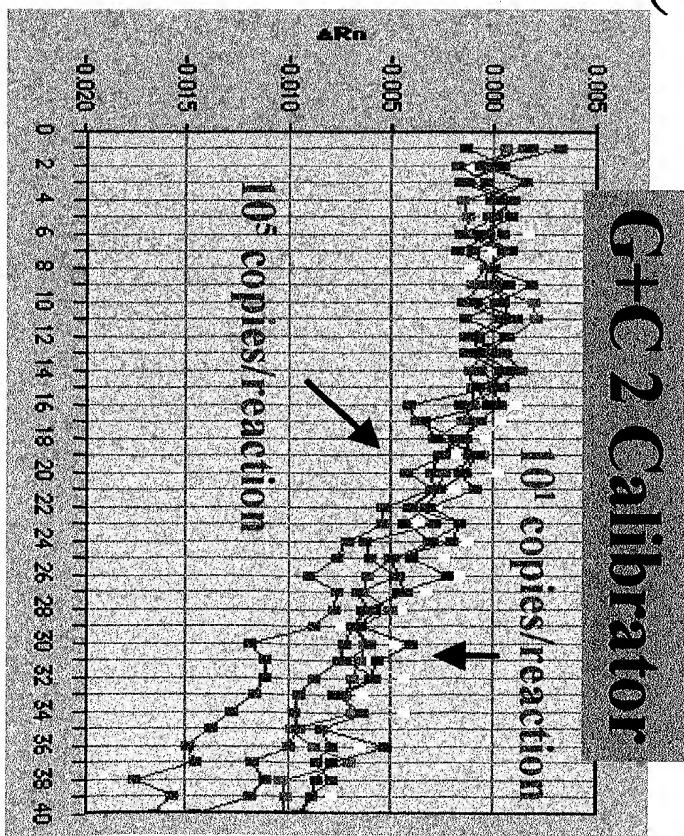


Table 1

HHV-7	Primers	Probes
	Forward (SEQ ID NO:1) 5'AGCGGTACCTGTAAATCATCCA3'	Standard (SEQ ID NO:3) 5' ACCAGTGAGAACATCGCTCTAACTGGATCA 3'
HHV-8	Reverse (SEQ ID NO:2) 5' AACAGAACGCCACCTCGAT 3'	Calibrator (SEQ ID NO:4) 5' TAAGCCCTGACCGCACGGGTATAACTAA 3'
	Primers	Probes
HHV-8	Forward (SEQ ID NO:5) 5'GTCCAGACGATATGTGCGC3'	Standard (SEQ ID NO:7) 5'CATTGGTGGTATATAGATCAAGTTCCGCCA3'
	Reverse (SEQ ID NO:6) 5'ACTCCAAAATATCGGCCGG3'	Calibrator (SEQ ID NO:8) 5'ACTATTCCATGCCGAATTCGAGCATAGTTG3'

(continued)

Table 1 (continued)

HIV-1	Primers	Probes
	Forward (SEQ ID NO:9) 5' TACTGACGCTCTCGCACC 3'	Standard (SEQ ID NO:11) 5' ATCTCTCTCTAGCCTCCGCTAGTCAA 3'
CAMV	Reverse (SEQ ID NO:10) 5' TCTCGACGCAGGGACTCG 3'	Calibrator (SEQ ID NO:12) 5' ACTCTCAGCGGCATTCTCCTCACTTCTACT 3'
	Primers	Probes
	Forward (SEQ ID NO:13) 5' GTCTTGCAGAGGATAGTGGGA 3'	Standard (SEQ ID NO:15) 5' TGCAGTCATCCCTTACGTCAGTGGAGAT 3'
	Reverse (SEQ ID NO:14) 5' CACGTTCAAGCAACTGGGA 3'	Calibrator (SEQ ID NO:16) 5' ATCGCTACATGCTAGGCATCTGTGTC 3'

(continued)

Table 1 (continued)

	Primers	Probes
Myc.	Forward (SEQ ID NO:17) 5' AGGAGGAGTGGCGCTGATG 3'	Standard (SEQ ID NO:19) 5' ACGAGGAGTCGCTGGCCCGATCC 3'
T. 1	Reverse (SEQ ID NO:18) 5' ACTCGGGCAGAGCTGCC 3'	Calibrator (SEQ ID NO:20) 5' TCCAGCGTCAGGCGTAGGCAGC 3'
	Primers	Probes
Myc.	Forward (SEQ ID NO:21) 5' AGGCGAACCCCTGCCAG 3'	Standard (SEQ ID NO:23) 5' TCGACACATAGGTGAGGTCTGCTACCCACA 3'
T. 2	Reverse (SEQ ID NO:22) 5' GATCGCTGATCCGCCA 3'	Calibrator (SEQ ID NO:24) 5' ACTACGACTACGGCTGCCATCGACATCGAT 3'

Table 2

Primers	Probes
Forward (SEQ ID NO:25) 5' CAAAGCCAAATTATCCAGAGCG 3'	Standard (SEQ ID NO:27) 5' CACCAGACGTCACACCCGAAGGAAT 3'
Reverse (SEQ ID NO:26) 5' CGCTAGGTTGAGGATCGA 3'	Calibrator (SEQ ID NO:28) 5' TACGGCAACGCCAACAGACCTAGCGA 3'
HHV-6	Calibrator
Primer forward (SEQ ID NO:29) 5' CCGGAAACCGAACATTACTGAA 3'	Probe (SEQ ID NO:31) 5' TACGGCAACGCCAACAGACCTAGCGA 3'
Primer reverse (SEQ ID NO:30) 5' TTACGTGAGGATTCGAGGGC 3'	

Table 1

	Primers	Probes
HHV-7		
Forward	<u>(SEQ ID NO:1)</u>	Standard <u>(SEQ ID NO:3)</u>
	5'AGCGGGTACCTGTAAAATCATCCA3'	5' ACCAGTGAGAACATCGCTCTAACTGGATCA 3'
Reverse	<u>(SEQ ID NO:2)</u>	Calibrator <u>(SEQ ID NO:4)</u>
	5' AACAGAACGCCACCTCGAT 3'	5' TAAGCCCTGACCGCACGGGTATAACTAA 3'
HHV-8		
Primers	Primers	Probes
Forward	<u>(SEQ ID NO:5)</u>	Standard <u>(SEQ ID NO:7)</u>
	5'GTCCAGACGATATGTGCGC3'	5'CATTGGTGGTATAGATCAAGTTCCGCCA3'
Reverse	<u>(SEQ ID NO:6)</u>	Calibrator <u>(SEQ ID NO:8)</u>
	5'ACTCCAAAATATCGGCCGG3'	5'ACTATTCCATGCGGAATTCGAGCATAGTTG3'

(continued)

Table 1 (continued)

	Primers	Probes
HIV-1		
Forward	<u>(SEQ ID NO:9)</u>	Standard <u>(SEQ ID NO:11)</u>
	5' TACTGACCGCTCTGGCAC 3'	5' ATCTCTCTCCTCTAGCCTCCGCTAGTCAA 3'
Reverse	<u>(SEQ ID NO:10)</u>	Calibrator <u>(SEQ ID NO:12)</u>
	5' TCTCGACGCCAGGACTCG 3'	5' ACTCTCAGCGGCATTCTCCTCACTTCTACT 3'
CAMV		
Primers		Probes
Forward	<u>(SEQ ID NO:13)</u>	Standard <u>(SEQ ID NO:15)</u>
	5' GTCTTGCGAAGGATAGTGGGA 3'	5' TGCCTCATCCCTTACGTCAGTGGAGAT 3'
Reverse	<u>(SEQ ID NO:14)</u>	Calibrator <u>(SEQ ID NO:16)</u>
	5' CACGTTCAAAGCAAGTGG 3'	5' ATCGCTACATGCTAGGCATCTGTGTGC 3'

(continued)

Table 1 (continued)

	Primers	Probes
Myc.		
T. 1	Forward <u>(SEQ ID NO:17)</u> 5' AGGAGGAGTGGCGCTGATG 3' Reverse <u>(SEQ ID NO:18)</u> 5' ACTCGGCCGAGAGCTGCC 3'	Standard <u>(SEQ ID NO:19)</u> 5' ACGAGGAGTGGCTGGCCGATCC 3' Calibrator <u>(SEQ ID NO:20)</u> 5' TCCAGCGTCAGGCGTAGGCAGC 3'
Myc.	Primers	Probes
T. 2	Forward <u>(SEQ ID NO:21)</u> 5' AGGCGAACCCCTGCCAG 3' Reverse <u>(SEQ ID NO:22)</u> 5' GATCGCTGATCCGGCCA 3'	Standard <u>(SEQ ID NO:23)</u> 5' TCGACACATAGGTGAGGTCTGCTACCCACA 3' Calibrator <u>(SEQ ID NO:24)</u> 5' ACTACGACTACGGCTGCGATCGACATCGAT 3'

Table 2

Primers	Probes
Forward (<u>SEQ ID NO: 25</u>) 5' CAAAGCCAAATTATCCAGAGCG 3'	Standard (<u>SEQ ID NO: 27</u>) 5' CACCAGACGGTCACACCCGAAGGAAT 3'
Reverse (<u>SEQ ID NO: 26</u>) 5' CGCTAGGTTGAGGATGATCGA 3'	Calibrator (<u>SEQ ID NO: 28</u>) 5' TACGCAACGCCAACAGACCTAGCGA 3'
HHV-6	Calibrator
Primer forward (<u>SEQ ID NO: 29</u>) 5' CCGGAAACCGAACATTACTGAA 3'	Probe (<u>SEQ ID NO: 31</u>) 5' TACGCCAACGCCAACAGACCTAGCGA 3'
Primer reverse (<u>SEQ ID NO: 30</u>) 5' TTACGTGAGGATCGAGGC 3'	